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      .      *::* ::* * .. . : . *****:*:* :*** *:::~ ***** * *:*..* :*:*:*:* . ** :~: :~ :*~ .
gi | 73974959 | ref | XP_855574.1 | MTSKLAVALLA AFVLSAALCEAAVLSRVSSSELRCQCIKTHSTPFHPKYIKELRVIDSGPHCENSEIIVKLFNGNEVCLDPKEKWWQKVVQIFLKKAEKQDP--- 101
gi | 27806863 | ref | NP_776350.1 | MTSKLAVALLA AFLLSAALCEAAVLSRMSTELRCQCIKTHSTPFHPKFIFELRVIESGPHCENSEIIVKLTNGNEVCLNPKEKWWQKVVQVFKRAEKQDP--- 101
gi | 10834978 | ref | NP_000575.1 | MTSKLAVALLA AFLIS AALCEGAVLPRSAKELRCQCIKTYSKPFHPKFIFELRVIESGPHCANTEIIVKLS DGRELCLDPKENWVQRVVEKFLKRAENS----- 99
gi | 114594298 | ref | XP_001156375.1 | MTSKLAVALLA AFLIS AALCEGAVLPRSAKELRCQCIKTYSKPFHPKFIFELRVIESGPHCANTEIIVKLS DGRELCLDPKENWVQRVVEKFLKRAENS----- 99
gi | 45384324 | ref | NP_990349.1 | MMGKAVAAVMALLLISMA GAKGMAQARS AIELRCQCIETHSKFIHPKFIQNVNLTSPGPHCKNVEVIATLKDGREVCLDPTAPWVKLIKAILDKADTNKKTAS 104
gi | 49169793 | ref | NP_990829.1 | -MNGKLGAVLALLLVSAALSQGRITLVKMGNELRCQCIETHSKFIHPKSIQDVKLTSPGPHCKNVEIATLKDGREVCLDPTAPWVQLIVKALMAKQLNSDAPL 103
1.....10.....20.....30.....40.....50.....60.....70.....80.....90.....100....

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